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On page 196, please delete the abstract on lines 2-18
and substitute therefor:

A2 The invention provides a method for identifying a polypeptide pharmacofamily. The method includes the steps of (a) determining bound conformations of a ligand bound to different polypeptides of a polypeptide family, and (b) identifying two or more bound conformations of the ligand having substantially different bound conformations, thereby identifying at least two polypeptide pharmacofamilies exhibiting binding specificity for the two or more substantially different bound conformations of the ligand.

In the claims:

Please cancel claims 1-18, 20, and 24-44.

Please amend the claims as follows:

19. A method for identifying polypeptide pharmacofamilies, comprising:

(a) determining bound conformations of a ligand bound to different polypeptides of a polypeptide family;

A3 (b) identifying two or more bound conformations of said ligand having substantially different bound conformations, thereby identifying at least a first bound conformation and a second bound conformation of said ligand;